



1600

**Does Not Comply
Corrected Diskette Needed**

RAW SEQUENCE LISTING

DATE: 09/11/2002

PATENT APPLICATION: US/09/392,822A

TIME: 09:06:09

Input Set : D:\Seqlist.txt

Output Set: N:\CRF4\09112002\I392822A.raw

4 <110> APPLICANT: Yu, De Chao
 5 Henderson, Daniel
 8 <120> TITLE OF INVENTION: Adenovirus Vectors Containing Cell
 9 Status-Specific Response Elements and Methods of Use Thereof
 12 <130> FILE REFERENCE: CELL-014; 122.1
 14 <140> CURRENT APPLICATION NUMBER: 09/392,822A
 15 <141> CURRENT FILING DATE: 1999-09-09
 17 <150> PRIOR APPLICATION NUMBER: 60/099,791
 18 <151> PRIOR FILING DATE: 1998-09-10
 20 <150> PRIOR APPLICATION NUMBER: 09/392,822
 21 <151> PRIOR FILING DATE: 1999-09-09
 23 <160> NUMBER OF SEQ ID NOS: 8
 25 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

60 <210> SEQ ID NO: 2
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 62 <212> TYPE: DNA
 63 <213> ORGANISM: H. sapiens
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 74 ttgactttta gtatccagcc cctctagaaa tctagctgat atagtgtggc tcaaaacctt 540
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 77 tggaaaagaa aaagaaagaa agggaaaaaa aaaaaaaa gagatgacct ctcaggctct 720
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 79 agggtcattg tgacgatcaa atgtggtcac gtgtatgagg caccagcaca tgcctggctc 840
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Insert hand return

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insert hand return

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TIME: 09:06:09

Input Set : D:\Seqlist.txt

Output Set: N:\CRF4\09112002\I392822A.raw

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/11/2002
PATENT APPLICATION: US/09/392,822A TIME: 09:06:10

Input Set : D:\Seqlist.txt
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Skipped Sequences(NEW RULES):

Sequence(s)___missing. If intentional, please use the following format for each skipped sequence.

<210> sequence id number

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Seq#:3,4,5

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Input Set : D:\Seqlist.txt

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L:163 M:254 E: No. of Bases conflict, LENGTH:Input:3 Counted:5838 SEQ:2
L:163 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:9
L:163 M:252 E: No. of Seq. differs, <211> LENGTH:Input:5836 Found:5838 SEQ:2
L:168 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:168 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:2 differs:3
M:254 Repeated in SeqNo=2
L:419 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:10
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L:424 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:424 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:2 differs:4
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L:625 M:252 E: No. of Seq. differs, <211> LENGTH:Input:12047 Found:12049 SEQ:0
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L:630 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:2 differs:5
L:647 M:216 E: (34) Seq.#s missing, SEQ ID NOS: 3 thru 5